

Appl. No. : Unknown
Filed : Herewith

(micro)organisms among other ones having homologous nucleotide sequences by identification of their nucleotide sequences, after amplification by a single primer pair.

Please replace the paragraph beginning at page 2, line 1, with the following rewritten paragraph:

Description of the Related Art

Please replace the paragraph beginning at page 4, line 28, with the following rewritten paragraph:

Summary of the Invention

Please replace the paragraph beginning at page 5, line 27, with the following rewritten paragraph:

Detailed Description of the Preferred Embodiments

Please replace the paragraph beginning at page 5, line 28, with the following rewritten paragraph:

The inventors have discovered that it is possible to drastically simplify the identification of one or several (micro)organisms among many other ones having homologous sequences by combining a single amplification using common primer pair and an identification of the possible (micro)organism(s) by detecting and possibly recording upon an array the presence of a single signal resulting only from binding between a capture sequence and its corresponding target sequence and correlating the presence of said detected target sequence to the identification of a genetic sequence specific of said (micro)organism(s). This means that the method and device according to the invention will allow the easy identification/detection of a specific sequence among other homologous sequences and its quantification (characterisation of the number of copies or presence of said organisms in a biological sample) of a target sequence, said target sequence having a nucleotide sequence specific of said (micro)organisms.

Please delete the paragraphs beginning at page 20, line 11 through the paragraph ending at page 20, line 27 and move them to page 5, line 12.

Please insert the paragraph at page 20 line 28 as follows:

Examples